SEQUENCE LISTING



<110> Skånemejerier AB

<120> NEW ENZYME AND ITS USE

<130> 75086

<150> US 60/320,139

<151> 2003-04-24

<150> US 60/481,598

<151> 2003-11-05

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<170> PatentIn version 3.2

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Val Asp Thr Pro Asn Leu Asp Ala Met Ala Arg Asp Gly Val Lys Ala 50 55 60

Arg Tyr Met Thr Pro Ala Phe Val Thr Met Thr Ser Pro Cys His Phe 65 70 75 80

Thr Leu Val Thr Gly Lys Tyr Ile Glu Asn His Gly Val Val His Asn 90 95

Met Tyr Tyr Asn Thr Thr Ser Lys Val Lys Leu Pro Tyr His Ala Thr 100 105 110

Leu Gly Ile Gln Arg Trp Trp Asp Asn Gly Ser Val Pro Ile Trp Ile 115 120 125

Thr Ala Gln Arg Gln Gly Leu Arg Ala Gly Ser Phe Phe Tyr Pro Gly

130 135 140

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Gly	Ile	Ala	His	Asn 165	Tyr	Lys	Asn	Glu	Thr 170	Glu	Trp	Arg	Ala	Asn 175	Ile
Asp	Thr	Val	Met 180	Ala	Trp	Phe	Thr	Glu 185	Glu	Asp	Leu	Asp	Leu 190	Val	Thr
Leu	Tyr	Phe 195	Gly	Glu	Pro	Asp	Ser 200	Thr	Gly	His	Arg	Tyr 205	Gly	Pro	Glu
Ser	Pro 210	Glu	Arg	Arg	Glu	Met 215	Val	Arg	Gln	Val	Asp 220	Arg	Thr	Val	Gly
Tyr 225	Leu	Arg	Glu	Ser	Ile 230	Ala	Arg	Asn	His	Leu 235	Thr	Asp	Arg	Leu	Asn 240
Leu	Ile	Ile	Thr	Ser 245	Asp	His	Gly		Thr 250	Thr	Val	Asp	Lys	Arg 255	
Gly	Asp	Leu	Val 260	Glu	Phe	His	Lys	Phe 265	·Pro	Asn	Phe	Thr	Phe 270	Arg	Asp
Ile	Glu	Phe 275	Glu	Leu	Leu	Asp	Tyr 280	Gly	Pro	Asn	Gly	Met 285	Leu	Leu	Pro
Lys	Glu 290	Gly	Arg	Leu	Glu	Lys 295	Val	Tyr	Asp	Ala	Leu 300	Lys	Asp	Ala	His
Pro 305	Lys	Leu	His	Val	Tyr 310	Lys	Lys	Glu	Ala	Phe 315	Pro	Glu	Ala	Phe	His 320
Tyr	Ala	Asn	Asn	Pro 325	Arg	Val	Thr	Pro	Leu 330	Leu	Met	Tyr	Ser	Asp 335	Leu
Gly	Tyr	Val	Ile 340	His	Gly	Arg	Ile	Asn 345	Val	Gln	Phe	Asn	Asn 350	Gly	Glu
His	Gly	Phe 355	Asp	Asn	Lys	Asp	Met 360	Asp	Met	Lys	Thr	Ile 365	Phe	Arg	Ala
Val	Gly 370	Pro	Ser	Phe	Arg	Ala 375	Gly	Leu	Glu	Val	Glu 380	Pro	Phe	Glu	Ser

Val His Val Tyr Glu Leu Met Cys Arg Leu Leu Gly Ile Val Pro Glu 385 390 395 400

Ala Asn Asp Gly His Leu Ala Thr Leu Leu Pro Met Leu His Thr Glu
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Ser Ala Leu Pro Pro Asp Ala Leu Leu Val Ala Asp Gly Pro Cys Leu 420 425 430

Pro Ser Leu Ser Gln Ala Lys Gly Cys Met Pro Leu Ser Pro Ala Ala 435 · 440 445

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gcatccagag gtggtgggac aacggcagcg tgcccatctg gatcacagcc cagaggcagg 420

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Val Asp Thr Pro Asn Leu Asp Ala Met Ala Arg Asp Gly Val Lys Ala 50 55 60

Arg Tyr Met Thr Pro Ala Phe Val Thr Met Thr Ser Pro Cys His Phe 65 70 75 80

Thr Leu Val Thr Gly Lys Tyr Ile Glu Asn His Gly Val Val His Asn 90 95

Met Tyr Tyr Asn Thr Thr Ser Lys Val Lys Leu Pro Tyr His Ala Thr 100 105 110

Leu Gly Ile Gln Arg Trp Trp Asp Asn Gly Ser Val Pro Ile Trp Ile 115 120 125

Thr Ala Gln Arg Gln Gly Leu Arg Ala Gly Ser Phe Phe Tyr Pro Gly 130 135 140

Gly 145	Asn	Val	Thr	Tyr	Gln 150	Gly	Val	Ala	Val	Thr 155	Arg	Ser	Arg	Lys	Glu 160
Gly	Ile	Ala	His	Asn 165	Tyr	Lys	Asn	Glu	Thr 170	Glu	Trp	Arg	Ala	Asn 175	Ile
Asp	Thr	Val	Met 180	Ala	Trp	Phe	Thr	Glu 185	Glu	Asp	Leu	Asp	Leu 190	Val	Thr
Leu	Tyr	Phe 195	Gly	Glu	Pro	Asp	Ser 200	Thr	Gly	His	Arg	Tyr 205	Gly	Pro	Glu
Ser	Pro 210	Glu	Arg	Arg	Glu	Met 215	Val	Arg	Gln	Val	Asp 220	Arg	Thr	Val	Gly
Tyr 225	Leu	Arg	Glu	Ser	Ile 230	Ala	Arg	Asn	His	Leu 235	Thr	Asp	Arg	Leu	Asn 240
Leu	Ile	Ile	Thr	Ser 245	Asp	His	Gly	Met	Thr 250	Thr	Val	Asp	Lys	Arg 255	Ala
Gly	Asp	Leu	Val 260	Glu	Phe	His	Lys	Phe 265	Pro	Asn	Phe	Thr	Phe 270	Arg	Asp
Ile	Glu	Phe 275	Glu	Leu	Leu	Asp	Tyr 280	Gly	Pro	Asn	Gly	Met 285	Leu	Leu	Pro
Lys	Glu 290	Gly	Arg	Leu	Glu	Lys 295	Val	Tyr	Asp	Ala	Leu 300	Lys	Asp	Ala	His
Pro 305	Lys	Leu	His	Val	Tyr 310	Lys	Lys	Glu	Ala	Phe 315	Pro	Glu	Ala	Phe	His 320
Tyr	Ala	Asn	Asn	Pro 325	Arg	Val	Thr	Pro	Leu 330	Leu	Met	Tyr	Ser	Asp 335	Leu
Gly	Tyr	Val	Ile 340	His	Gly	Arg	Ile	Asn 345	Val	Gln	Phe	Asn	Asn 350	Gly	Glu
His	Gly	Phe 355	Asp	Asn	Lys	Asp	Met 360		Met	Lys	Thr	Ile 365	Phe	Arg	Ala
Val	Gly 370	Pro	Ser	Phe	Arg	Ala 375	Gly	Leu	Glu	Val	Glu 380	Pro	Phe	Glu	Ser
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385 390 395 400

Ala Asn Asp Gly His Leu Ala Thr Leu Leu Pro Met Leu His Thr Glu 405 410 415

Ser Ala Leu Pro Pro Asp Gly Arg Pro Thr Leu Leu Pro Lys Gly Arg 420 425 430

Ser Ala Leu Pro Pro Ser Ser Arg Pro Leu Leu Val Met Gly Leu Leu 435 440 445

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gcatccagag gtggtgggac aacggcagcg tgcccatctg gatcacagcc cagaggcagg 420

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Leu Leu Val Ser Phe Asp Gly Phe Arg Trp Asn Tyr Asp Gln Asp 35 40 45

Val Asp Thr Pro Asn Leu Asp Ala Met Ala Arg Asp Gly Val Lys Ala 50 55 60

Arg Tyr Met Thr Pro Ala Phe Val Thr Met Thr Ser Pro Cys His Phe 65 70 75 80

Thr Leu Val Thr Gly Lys Tyr Ile Glu Asn His Gly Val Val His Asn 90 95

Met Tyr Tyr Asn Thr Thr Ser Lys Val Lys Leu Pro Tyr His Ala Thr 100 105 110

Leu Gly Ile Gln Arg Trp Trp Asp Asn Gly Ser Val Pro Ile Trp Ile 115 120 125

Thr Ala Gln Arg Gln Gly Leu Arg Ala Gly Ser Phe Phe Tyr Pro Gly 130 135 140

Gly Asn Val Thr Tyr Gln Gly Val Ala Val Thr Arg Ser Arg Lys Glu
145 150 155 160

Gly Ile Ala His Asn Tyr Lys Asn Glu Thr Glu Trp Arg Ala Asn Ile

165 170 175

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Ser	Pro 210	Glu	Arg	Arg	Glu	Met 215	Val	Arg	Gln	Val	Asp 220	Arg	Thr	Val	Gly
Tyr 225	Leu	Arg	Glu	Ser	Ile 230	Ala	Arg	Asn	His	Leu 235	Thr	Asp	Arg	Leu	Asn 240
Leu	Ile	Ile	Thr	Ser 245	Asp	His	Gly	Met	Thr 250	Thr	Val	Asp.	Lys	Arg 255	Ala
Gly	Asp	Leu	Val 260	Glu	Phe	His	Lys	Phe 265	Pro	Asn	Phe	Thr	Phe 270	Arg	Asp
Ile	Glu	Phe 275	Glu	Leu	Leu	Asp	Tyr 280	Gly	Pro	Asn	Gly	Met 285	Leu	Leu	Pro
Lys	Glu 290	Gly	Arg	Leu	Glu	Lys 295	Val	Tyr	Asp	Ala	Leu 300	Lys	Asp	Ala	His
Pro 305	Lys	Leu	His	Val	Tyr 310	Lys	Lys	Glu	Ala	Phe 315	Pro	Glu	Ala	Phe	His
Tyr	Ala	Asn	Asn	Pro 325	Arg	Val	Thr	Pro	Leu 330	Leu	Met	Tyr	Ser	Asp 335	Leu
Gly	Tyr	Val		His							Phe		Asn 350		Glu
His	Gly	Phe 355	Asp	Asn	Lys	Asp	Met 360	Asp	Met	Lys	Thr	Ile 365	Phe	Arg	Ala
Val	Gly 370	Pro	Ser	Phe	Arg	Ala 375	Gly	Leu	Glu	Val	Glu 380	Pro	Phe	Glu	Ser
Val 385	His	Val	Tyr	Glu	Leu 390	Met	Cys	Arg	Leu	Leu 395	Gly	Ile	Val	Pro	Glu 400
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